5

10

15

20

25

30

What Is Claimed Is:

1. A recombinant Equine Herpes Virus (EHV) wherein the protein gM is absent, and wherein said EHV is free of heterologous elements.

- 2. The EHV according to claim 1, wherein the gene coding for the protein gM is deleted.
- 3. The EHV according to claim 1, obtainable by a method comprising the steps of:
 - a) isolating a wild-type EHV;
 - establishing a plasmid encoding the EHV gM gene, optionally with flanking sequences;
 - c) generating a complementing cell line expressing gM or parts thereof;
 - d) establishing an EH virus carrying a GFP-encoding cassette insert in its gM coding sequence by co-transfecting the complementing cell line of step b) with EHV-nucleic acid and a plasmid encoding gM which is interrupted by a GFP-encoding cassette insert;
 - e) deleting the GFP-encoding cassette; and
 - selecting for the EHV clones wherein the GFP-encoding cassette is successfully deleted.
- 4. The EHV according to claim 1, wherein the EHV is EHV-1.
- 5. The EHV-1 according to claim 4, wherein 850-1100 bp of the gM open reading frame are deleted.
- 6. The EHV-1 according to claim 4, wherein the entire gM coding sequence is deleted except for 150-200 bp of the coding sequence for the C-terminal portion and except for 150-250 bp of the coding sequence for the N-terminal portion.
- 7. The EHV-1 according to claim 6, wherein the entire gM coding sequence is deleted except for 184 bp of the coding sequence for the C-terminal portion and except for 209 bp of the coding sequence for the N-terminal portion.

10

25

30

8. The EHV-1 according to claim 4, wherein nucleotides 93268-93318 to 94222-94322 of the gM coding sequence as corresponding to SEQ ID NO:1 are deleted.

- The EHV-1 according to claim 7, wherein nucleotides 93268 to 94322 of the gM coding sequence as corresponding to SEQ ID NO:1 are deleted.
 - 10. The EHV-1 according to claim 8, wherein nucleotides 94263 to 93302 of the gM coding sequence as corresponding to SEQ ID NO:1 are deleted.
 - 11. The EHV-1 according to claim 4, wherein said EHV-1 is a recombinant variant based on strain RacH of EHV-1.
- 12. The EHV-1 according to claim 11, wherein said EHV-1 is RacH-based recombinant variant isolate HΔgM-w as deposited at the ECACC/CAMR on October 16, 2002 with the accession number 02101663.
 - 13. The EHV according to claim 1, wherein said EHV is EHV-4.
- ²⁰ 14. The EHV-4 according to claim 13, wherein 900-1150 bp of the gM open reading frame are deleted.
 - 15. The EHV-4 according to claim 13, wherein the entire gM coding sequence is deleted except for 0-50 bp of the coding sequence for the C-terminal portion and except for 150-250 bp of the coding sequence for the N-terminal portion.
 - 16. The EHV-4 according to claim 15, wherein the entire gM coding sequence is deleted except for 34 bp of the coding sequence for the C-terminal portion and except for 209 bp of the coding sequence for the N-terminal portion.
 - 17. The EHV-4 according to claim 13, wherein nucleotides 92681-92731 to 93765-93865 of the gM coding sequence as corresponding to SEQ ID NO:2 are deleted.

15

25

30

18. The EHV-4 according to claim 16, wherein nucleotides 92681 to 93865 of the gM coding sequence as corresponding to SEQ ID NO:2 are deleted.

- 19. The EHV-4 according to claim 17, wherein nucleotides 92715 to 93824 of the gM coding sequence as corresponding to SEQ ID NO:2 are deleted.
- 20. The EHV-4 according to claim 13, wherein said EHV-4 is a recombinant variant based on MSV Lot 071398 of EHV-4.
- 21. The EHV-4 according to claim 20, wherein said EHV-4 is based on MSV Lot 071398 and isolate E4ΔgM-w and that it is the EHV-4 which was deposited at the ECACC/CAMR on January 14, 2003 with the accession number 03011401.
 - 22. A nucleic acid coding for an EHV according to claim 1.
 - 23. A vaccine preparation comprising an EHV according to claim 1.
 - 24. A vaccine preparation comprising the nucleic acid according to claim 22.
- 25. A vaccine preparation comprising at least one EHV-1 according to claim 4 and one EHV-4 according to claim 13.
 - 26.A method of treatment and/or prevention of an EHV-associated condition comprising administering to a mammal in need of such a treatment a therapeutically effective amount of the vaccine preparation according to claim 23 or 24 and monitoring the therapeutic success.
 - 27.A method of treatment and/or prevention of an EHV-associated condition comprising administering to a mammal in need of such a treatment a therapeutically effective amount of the vaccine preparation according to claim 25 and monitoring the therapeutic success.
 - 28. A method for obtaining a recombinant EHV, comprising the steps of:

10

15

- a) isolating a wild-type EHV;
- b) establishing a plasmid encoding the EHV gM gene, optionally with flanking sequences;
- c) generating a complementing cell line expressing gM or parts thereof;
- d) establishing an EH virus carrying a GFP-encoding cassette insert in its gM coding sequence by co-transfecting the complementing cell line of step b) with EHV-nucleic acid and a plasmid encoding gM which is interrupted by a GFP-encoding cassette insert;
- e) deleting the GFP-encoding cassette; and
- f) selecting for the EHV clones wherein the GFP-encoding cassette is successfully deleted.
- 29. A cell line for use in a method according to claim 28, wherein the gene encoding the protein gM is transfected into said cell line and said cell line expresses gM.
- 30. The cell line according to claim 29, wherein the cell line is a cell line selected from the group of Vero cells, RK-13, and cc.
- 31. The cell line according to claim 30, wherein the cell line is the gM complementing cell line VERO GM which was deposited at the ECACC/CAMR on January 28, 2003 with the accession number 03012801.